Run 3

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Result.
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062640 sus scrofa
Q13490 homo sapien
Q30660 gallus gall
Q30660 gallus gall
Q30660 gallus gall
Q30660 gallus gall
Q30210 mus musculu
Q41437 orgyla pomon
Q91166 mus musculu
Q99485 mus musculu
Q99485 mus musculu
Q99485 mus musculu
Q99486 grattus norv
P98170 homo sapien
Q4307 drosophila
Q34307 drosophila
Q24307 drosophila
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BIR2_HUMAN
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45	44	43	42	41	40	39	38	37	36	35	34	
54	54.5	54.5	54.5	55	56	56.5	56.5	56.5	57	58.5	58.5	
18.4	18.5	18.5	18.5	18.7	19.0	19.2	19.2	19.2	19.4	19.9	19.9	
64	1170	1170	1170	361	65	249	156	114	1004	1173	844	
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MTA_STRPU	TSP1_MOUSE	TSP1_HUMAN	TSP1_BOVIN	VPI_BPMU	MTB_STRPU	IAP2_NPVAC	BODP_HUMAN	IBB2_PEA	PO11_SCICO	TSP1_XENLA	PBPB_ECOLI	
P04734 strongyloce	P35441 mus musculu	P07996 homo sapien	Q28178 bos taurus	Q01267 bacteriopha	Q27287 strongyloce	P41454 autographa	P36639 homo sapien	Q41066 pisum sativ	Q03277 sciara copr		P02919 e penicilli	

ALIGNMENTS

Oliabs, olicage, olivarian, oliabs, oliabs, oliabs, olicabe, oliabs, o	FORM AN HETEKOMENIC COMPLEX, WHICH IS THEM RECRUITED TO THE TUMOR NECROSIS FACTOR RECEPTOR 2 (THEFE). -(- SUBCELULIAR LOCATION: CYROPLASHIC (POTENTIAL).	888
01489; 016628; 091946; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 01-NOV-1997 (Rel. 35, Last sequence update) 20-AUG-2001 (Rel. 40, Last annotation update) 20-AUG-2001 (Augustion upda	-i- FUNCTION: APOPTOTIC SUPPRESSOR. THE BIR MOTIFS REGION I: WITH THE RECEPTOR ASSOCIATED FACTORS 1 AND 2 (TRAFT AND	388
OliABS; OliCare Summary, OliCare Summary, OliABS; OliCare Summary, OliCar		23:
OliABS; OliCoza; Schwark, OliABS; OliCoza; OliABS; O	<pre>including inhibitor of apoptosis protein=1, stannin, and two novel</pre>	7 P.
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OliABS; OliCoza; Surmark, OliABS; OliCoza; Rel. 35, Created) OliABS; OliCoza; OliABS; OliCoza; OliABS; OliCoza; OliABS; OliCoza; Rel. 35, Last sequence update) OliAOS-1997 (Rel. 35, Last sequence update) OliAOS-1997 (Rel. 35, Last sequence update) OliAOS-2001 (Rel. 40, Last annotation update) OliA	A	RX.
OliABS; OliAG2B; OlPH46; OliABS; OliCord Rel. 35, Created) Ol-NOV-1997 (Rel. 35, Created) Ol-NOV-1997 (Rel. 35, Last sequence update) Ol-NOV-1997 (Rel. 35, Created) BACULOVIRAL IAP REPEAT-COVYANING PROTEIN 1 (IHHIBITOR OF APOP PROTEIN 1) (HAP1) (HIAP-1) (C-IAP2) (TNFR2-TRAF SIGNALING COM PROTEIN 1) (HAP1) (HIAP-1) (C-IAP2) (TNFR2-TRAF SIGNALING COM PROTEIN 1) (IAP HONGLOG C). BIRCE OR API1 OR IAP1 OR MIHC. HONG Sapiens (Human). BUKARYOLD: Wetzoar; Chordata; Craniata; Vertebrata; Euteleosto Mammalia; Eutheria; Pubmed-8548810; REDLINE-961201; Pubmed-8548810; REDLINE-961201; Pubmed-8548810; MEDLINE-961201; Pubmed-8548810; MEDLINE-961201; Pubmed-8548810; MEDLINE-961201; Pubmed-8552191; MEDLINE-96140249; Pubmed-8552191; MEDLINE-96140249; Pubmed-8552191; MEDLINE-96140249; Pubmed-8552191; MEDLINE-96100 of apoptosis in mammalian cells by NAIP and a rel Suppression of apoptosis inhibitory protein homolo function to inhibit apoptosis and/or bind tumor necrosis facto Proc. Natl. Acad. Sci. U.S. A. 93:4974-4978(1996).	SEQUENCE FROM N.A.	RP :
OliABS; OliAGZB; ODHAGA; OliABS; OliCoza, Created) Ol-NOV-1997 (Rel. 35, Created) Ol-NOV-1997 (Rel. 35, Last agquence update) Ol-NOV-1997 (Rel. 35, Last sequence update) Ol-NOV-1997 (Rel. 35, Created) BACULOVIRAL IAP REPEAT-COVYAINING PROTEIN 1 (IHHIBITOR OF APOP PROTEIN 1) (HAP1) (HIAP-1) (C-IAP2) (TNER2-TRAF SIGNALING COMPROTEIN 1) (IAP HONGLOG C). BIRCZ OR API1 OR IAP1 OR MIHC. HOMO Sapiens (Human) Dukaryoto: Metazoa; Chordata; Craniata; Vertebrata; Euteleosto Nemmalia; Eutheria; Primates; Catarrhini; Hominidme; Homo. II] ENDATA-96128127: Pubmed-8548810; MEDILINE-96128127: Pubmed-8548810; MEDILINE-96128127: Pubmed-8548810; MEDILINE-96128127: Pubmed-8548810; MEDILINE-9614924; Pubmed-8552191; MEDILINE-9614924; Pubmed-8552191; MEDILINE-9614924; Pubmed-8552191; MEDILINE-9614924; Pubmed-8552191; Nature 379:349-353(1996). SEQUENCE FROM N.A. TICSUE-Fetal liver; MEDILINE-9620843; Pubmed-8643514; Uren A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TICSUE-Fetal liver; MEDILINE-9520843; Pubmed-8643514; Uren A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D.L.; TUCEN A.G., Pakusah M., Hawkins C.J., Puls K.L., Vaux D	c. Natl. Acad. Sci. U.S.A.	R F
OliAND; OliAGZB; ODUPAG; OliAND; OliAGZB; ODUPAG; OliAND; OliAGZB; ODUPAG; OliAND; OliAGZB; ODUPAG; OliAND; OliAGZB; ORDIAG; OliAND; OliAGZB; ORDIAG; OliAND; OliAGZB; ORDIAG; OliAND; OliAGZB; ORDIAG; OliAND; OliAGZB; OliAGZB; OliAND; OliAGZB; OliAND; OliAGZB; OliAND; OliAGZB; OliAG	receptor-associated factors,";	27
Oli489; Oli6628; Opur46; Oli489; Oli6628; Opur46; Oli489; Oli6628; Opur46; Oli489; Oli628; Opur46; Opur48; Opur48; Opur48; Opur48; Oli489; Oli489; Opur48; Opu	function to inhibit apoptosis and/or bind tumor necrosis factor	RT
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OliABS; OliAG2B; ODUP46; OliABS; OliAG2B; ODUP46; OliAG9; OliAG797 (Rel. 35, Last sequence update) OliAG9-1097 (Rel. 35, Last sequence update) OliAG9-2001 (Rel. 40, Last annotation update) OliAG9-2001 (Rel. 40, Last annotation update) DO-AIG-2001 (Rel. 40, Last annotation update) PROTEIN 1) (HIAP1) (HIAP-1) (C-IAP2) (TWRZ-TRAF SIGNALING COMPROTEIN 1) (HIAP1) (HIAP-1) (C-IAP2) (TWRZ-TRAF SIGNALING COMPROTEIN 1) (HIAP1) (HIAP-1) (C-IAP2) (TWRZ-TRAF SIGNALING COMPROTEIN 1) (HIAP1) (HIAP1) (HIAP2) (TWRZ-TRAF SIGNALING COMPROTEIN 1) (HIAP1) (HIAP1) (HIAP2) (TWRZ-TRAF SIGNALING COMPROTEIN 1; HOMINIdae; HOMO. REBUZING-1812127; PubMed-8548810; RECLIR M., Pan M. G., Henzel W.J., Ayres T.M., Goeddel D.V.; FIDE TWRZ-TRAF Signaling complex contains two novel proteins to baculoviral inhibitor of apoptosis proteins. "; Cell 83:1243-1252(1995). INSUDEALIVET; MEDLINE-96149249; PubMed-8552191; Liston P., Roy Lean M., Ikeda J., Mackenzie A., Korneluk R.G.; "Suppression of apoptosis in mammalian cells by NAIP and a rel family of IAP genes."; Nature 379:349-353(1996). INSUDE-Fetal liver;	209843; Pubmed=8643514;	7 X
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Dirachoram Oli488; Oli6628; Opupu6; Oli489; Oli6628; Opupu6; Oli489; Oli6628; Opupu6; Oli489; Oli628; Opupu6; Oli489; Oli628; Opupu6; Oli489; Oli628; Opupu6; Oli489; Oli628; Opupu6; Oli489; Oli6828; Opupu6; Oli489; Oli489; Opupu6; Oli489; Oli489; Opupu6; Oli489; Ol		RP
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Oli489; Oli6628; Opud46; Oli489; Oli6628; Opud46; Oli489; Oli6628; Opud46; Oli489; Oli6628; Opud46; Oliky-1997 (Rel. 35, Last annotation update) Oliky-1997 (Rel. 35, Last annotation update) Oli697 (Rel. 35, Last annotation update) Oli697 (Rel. 40, Last annotation update) Oli697 (Rel. 41, Last annotation update) Oli697 (Independent update) Oli697 (Independen		5 R
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D1489; 016628; 020046; 011489; 016628; 020046; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 02-AIG-2001 (Rel. 40, Last annotation update) 03-AIG-2001 (Rel. 40, Last annotation update) 04-AIG-2001 (Rel. 40, Last annotation update) 04-AIG-2001 (Rel. 40, Last annotation update) 04-AIG-2001 (Rel. 40, Last annotation update) 05-AIG-2001 (Rel. 40, Last annotation update) 06-AIG-2001 (AIAP-1) (C-IAP2) (TNP72-TRAF SIGNALING COMPROTEIN 1) (IAPP 10, LAPP 1) (C-IAP2) (TNP72-TRAF SIGNALING COMPROTEIN 1) (IAPP 10, LAPP 1) (C-IAP2) (TNP72-TRAF SIGNALING CATAINTHIN; Hominidae; Homo. 05-AIG-1200-1200 (AIAP-1) (AIAP	Farahani R., McLean M., Ikeda J., Mackenzie A., Korneluk R.G.;	RA 3
Oli489; Oli6628; Olp46; Oli489; Oli689; Ole48; Ole48; Ole48; Oli489; Oli689; O	him o deind o oboston-Bosses	J X
Oli489; Ol6628; OgDP46; Oli489; Olf6628; OgDP46; Oli489; Olf628; Olf62		
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D1489; 016628; 020046; 011489; 016628; 020046; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 20-AUG-2001 (Rel. 40, Last annotation update) 20-AUG-2001 (Rel. 40, Last annotation) 20-AUG-2001 (Re		RN
Oli489; Ol6628; OgD466; Oli489; Ol6628; OgD466; Oli489; Olf628; OgD46; Oli489; Olf628; Olf	Cell B3:1243-1252(1995).	Z :
DIABN: 016628; 020046; 011489; 016628; 020046; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 01-NOV-1997 (Rel. 35, Last sequence update) 20-AIG-2001 (Rel. 40, Last annotation update) 20-AIG-2001 (ART annotation update) 20	Firtagord	2 2
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	Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Nammalia: Futboria: Drimates: Catarrhini: Hominidae	3 8
	Homo sapiens (Human).	So
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Pfam: PF00653; CARD: 1.
Pfam: PF00059; zf-C3HC4; 1.
SMARF; SM0023B; BIR; 3.
SMARF; SM00144; CARD; 1.
SMARF; SM00144; RING; 1.
                                                                                                              PIRP_PIG STANDARD,
062640;
052640;
15-DEC-1998 (Rel. 37, Created)
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ZN_FING
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InterPro; IPR001315; CARD.
InterPro; IPR001841; Znf_ring.
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SEQUENCE FROM N.A.
                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                Sus scrofa (Pig)
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REPEAT
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                                    NCBI_TaxID-9823;
                                                                                                                                                                                                                                                                                                                                          273
                                                                                                                                                                                                                                                                                                                                                             1 PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFFRCE 48
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SIMILARITY: CONTAINS 3 BIR REPEATS.
SIMILARITY: CONTAINS 1 CARD DOMAIN.
SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER
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llarity 100.0%;
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119
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163
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191
                                                         Chordata; Craniata; Vertebrata; Cetartiodactyla; Suina; Suidae;
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H -> P (IN
K -> P (IN
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Pred. No. 1.9
0; Mismatches
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-> P (IN REF. 2).
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ss 0;
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                                                                               Euteleostomi;
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Best Local
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Pfam; PF00653; BIR; 2.
Pfam; PF00653; CARD; 1.
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00218; BIR; 2.
SMART; SM00114; CARD; 1.
SMART; SM00114; CARD; 1.
SMART; SM00184; RING; 1.
PROSITE; PS01282; BIR_REPEAT_1: 2
PROSITE; PS50143; BIR_REPEAT_2: 2
PROSITE; PS50143; BIR_REPEAT_2: 2
PROSITE; PS50143; BIR_REPEAT_2: 2
                                                                                                                                                                                                                                                                                                                                                                         BIR3_HUMAN STANDARD; PRT; 618 AA.
013490; O16516;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
20-NOG-9001 (Rel. 40, Last annotation update)
BACULOVIRAL IAP HEPEAT-CONTAINING PROTEIN 3 (INHIBITOR OF APOPTOSIS PROTEIN 2) (HIAP-2) (HIAP-2) (C-IAPI) (TNFK2-TRAE SIGNALING COMPLEX PROTEIN 2) (JAP HONDIGG B).
BROTGEN 2) (JAP HONDIGG B).
BIRCS OR APIZ OR IAPZ OR MIHB.
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"Cytokine induced expression of porcine inhibitor of protein (iap) Kamily member is regulated by NF kappa Blochem, Biophys, Res. Commun. 243:827-832(1998).

- I. SIMILARITY: BOLONGS TO THE IAP FAMILY.

- I. SIMILARITY: CONTAINS 2 BIR REPEATS.

- I. SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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                                                       "The TNFR2-TRAF signalin to baculoviral inhibitor Cell 83:1243-1252(1995).
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ZN_FING
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Stehlik C., de Martin R., Binder
                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
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TISSUE-Liver;
                          SEQUENCE FROM N.A
                                                                                                     Rothe M., Pan M.-G., Hanzel W.J., Ayres T.M., Goeddel D.V., Rothe M., Pan M.-G., Hanzel W.J., Ayres T.M., Goeddel D.V., The Third That is two novel proteins obaculoviral inhibitor of apoptosis proteins.";
                                                                                                                                                                                           SEQUENCE FROM N.A. MEDLINE-96128127; PubMed-8548810;
                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR001370; BIR. InterPro; IPR001315; CARD.
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                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
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45; Conser
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4 70 BJ
90 157 BJ
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311
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93.8%;
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Pred. No. 2.5e
3; Mismatches
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RING-TYPE.
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Catarrhini;
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Pfam; PF00619; CARD; 1.
Pfam; PF009097; zf-C3HC4; 1.
SMART; SM00238; BIR; 3.
SMART; SM00184; RING; 1.
SMART; SM00184; RING; 1.
SMART; SM00184; RING; 1.
                       CONFLICT
CONFLICT
CONFLICT
                                                                                                                          REPEAT
DOMAIN
ZN_FING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; L49431; AAC41942.1; -.
EMBL; U45879; AAC50372.1; -.
EMBL; U37547; AAC50508.1; -.
PDB; 1QBH; 20-OCT-99.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SMISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in oway modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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Nat. St
-i- FUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Uren A.G., Pakusch M., Hawkins C.J., Puls K.L., "Cloning and expression of apoptosis inhibitory function to inhibit apoptosis and/or bind tumor receptor associated factors.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Liston P., Roy N., Tamai K., Lefebvre C., Baird Farahani R., McLean M., Ikeda J., Mackenzie A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE-Fetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nature 379:349-353(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Suppression of apoptosis in mammalian cells by family of IAP genes.";
                                                                                                                                                                                                                            Apoptosis;
REPEAT
                                                                                                                                                                                                                                                                         PROSITE; PS50143; BIR_REPEAT_2; PROSITE; PS50209; CARD; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PDB; 1QBH; 2
MIM; 601721;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hinds M.G., Norton R.S., Vaux D.L., Day C.L.;
"Solution structure of a baculoviral inhibitor of apoptosis (IAP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE-99332054; PubMed-10404221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRUCTURE BY NMR OF 266-363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=96209843; PubMed=8643514;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=96149249; PubMed=8552191
                                                                                                                                                                                                    REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Struct. Biol. 6:648-651(1999).

Struct. Biol. 6:648-651(1999).

FUNCTION: APOPTOTIC SUPPRESSOR. THE BIR MOTIFS REGION INTERACTS FUNCTION: APOPTOTIC SUPPRESSOR. THE BIR MOTIFS REGION INTERACTS WITH THE RECEPTOR ASSOCIATED FACTORS I THEN RECRUITED TO THE TUMOR POORM AN HETEROMBERIC COMPLEX, WHICH IS THEN RECRUITED TO THE TUMOR NECROSIS FRACTOR RECEPTOR 2 (THER2).

SUBCELLULAR LOCATION: CYTOPLASMIC (POTENTIAL) AND ADULT TISSUES. TISSUE SPECIFICITY: PRESENT IN MANY FETAL AND ADULT TISSUES. MAINLY EXPRESSED IN ADULT SKELETAL MUSCLE. THYMUS, TESTIS, OVARY, AND PANCREAS, LOW OR ABSENT IN BRAIN AND PERIPHERAL BLOOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LEUKOCYTES
SIMILARITY: BELONGS TO THE TAP FAMILY.
SIMILARITY: CONTAINS 3 BIR REPEATS.
SIMILARITY: CONTAINS 1 CARD DOMAIN.
SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Natl. Acad. Sci. U.S.A. 93:4974-4978(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IPR001370; BIR.
IPR001315; CARD.
IPR001841; Znf_ring.
                                                                                                                                                                                                                                                    Zinc-finger;
                             184
269
453
571
157
308
414
514
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                                                  113
250
336
539
605
157
157
     69899
                                                                                                                                                                                                                                                    Repeat;
  ₹
:
                       BIR 1.
BIR 2.
BIR 3.
CARD.
RING-TYPE.
S -> P (IN
C -> P (IN
C -> W (IN
S -> P (IN REF. 2).
C -> G (IN REF. 2).
Q -> L (IN REF. 2).
L -> W (IN REF. 2).
C1778D328063586D (
                                                                                                                                                                                                                                                 3D-structure.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaux
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Korneluk R.G.;
     CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and a related
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HESOLUT 4

BIR_CHICK
ID BIR_C

AC 09060
DT 01-W0
DT 01-W0
DT 10-W0
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Best Local s
Matches 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BIR_CHICK
Q90660;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lymphocytes.";

DNA Ceil Biol. 15:981-988(1996).

- FUNCTION: APODTOTIC SUPPRESSOR (BY SIMILARITY).

- FUNCTION: APODTOTIC SUPPRESSOR (BY SIMILARITY).

- SUBCELLULAR LOCATION: PREDOMINANTLY MUCLEAR.

- TISSUE SPECIFICITY: CELLS OF THE TLYMHOCYTE LINEAGE. FOUND BOTH CORTICAL AND MEDULLARY CELLS OF THE THYMUS.

- DEVELOPMENTAL STAGE: HIGH LEVELS ARE INDUCED WITHIN 4-8 HOURS T-CELL ACTIVATION IN SPLEEM AND THYMUS.

- SIMILARITY: BELONGS TO THE IAP FAMILY.

- SIMILARITY: CONTAINS 3 BIR REPEATS.

- SIMILARITY: CONTAINS 3 BIR REPEATS.

- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gallus gallus (Chicken).
Eukaryota; Metazoa; Chor
                                                                             Apoptosis;
REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Digby M.R., Kimpton W.G., York J.J., Connick T.E., Lowenthal J.W.; "ITA, a vertebrate homologue of IAP that is expressed in T
                        REPEAT
ZN_FING
                                                                                                                                                                                   PROSITE;
                                                                                                                                                                                                                                  Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00238; BIR; 3.
SMART; SM00114; CARD; 1.
SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INHIBITOR OF APOPTOSIS PROTEIN (IAP) (INHIBITOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=97101112; PubMed-8945639;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Archosauria;
                                                                                                                                                           PROSITE;
                                                                                                                                                                                                                                                                                                                                        Pfam; PF00653; BIR; Pfam; PF00619; CARD;
                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR001370; BIR.
InterPro; IPR001315; CARD.
InterPro; IPR001841; Znf_r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; U27466; AAB48118.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45;
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                                                                                                                                                              PS50209; CARD;
                                                                                                                                                                                                            PS01282; BIR_REPEAT_1;
                                                                                                                                  Zinc-finger;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (Rel.
(Rel.
(Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                etazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           35,
40,
                                                                                                                                                                                   BIR_REPEAT_2:
      69009
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                                                                                                                                                                                                                                                                                                                                                                                             Znf_ring
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Last annotation update)
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                                                    Repeat; Nuclear pair 1.
BIR 1.
BIR 2.
BIR 3.
      MW;
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                               RING-TYPE
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      53FC9136F34EBDDD CRC64;
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5.5e-27;
                                                                                                                                     protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Š
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and for commercial
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STATES OF STATES
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                     Query Match
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DIRZ_MOUSE STANDARD; PRT; 600 AA.

008863;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
20-ANG-2001 (Rel. 40, Last annotation update)
BACULOVIRAL IAP REPERT-CONTAINING PROTEIN 2 (INHIBITOR OF APOPTOSIS
                                                                                      ZN_FING
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                              Pfam; F
Pfam; F
SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 between
the Euro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDILIRE-98110590; PubMed-9441758;
Liston P., Lefebvre C., Fong W.G., Xuan J.Y., Korneluk R.G.;
"Genomic characterization of the mouse inhibitor of apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
TISSUE-Skeletal muscle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROTEIN 1) (MIAP1) (MIAP-1).
BIRC2 OR API1 OR IAP1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOUSE
                                                                                                                                                                                                                                               PROSITE; PS01282; BIR_REPEAT_1; PROSITE; PS50143; BIR_REPEAT_2; PROSITE; PS50209; CARD; 1.
                                                                                                                                                                                                                                                                                                                           SMART
                                                                                                                                                                                                                                                                                                                                        SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  MGD; MGI:1197007; Birc2.
InterPro; IPR001370; BIR.
InterPro; IPR001315; CARD.
InterPro; IPR001841; Znf_r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; U88908; AAC53531.1; -.
                                                                                                                                                                                   REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genomics 46:495-503(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID-10090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: APOPROTIC SUPPRESSOR, THE BIR MOTIFS REGION INTERACTS
WITH TWE RECEPTOR ASSOCIATED FACTORS 1 AND 2 (TRAF1 AND TRAF2) TO
FORM AN HETEROMERIC COMPLEX, WHICH IS THEN RECRUITED TO THE TUMOR
RECROSIS FACTOR RECEPTOR 2 (TMFR2) (BY SIMILARITY)
SUBCELLULAR LOCATION: CYTOPLASMIC (POTENTIAL).
SIMILARITY: BELONGS TO THE IAP FAMILY.
SIMILARITY: CONTAINS 3 BIR REPEATS.
SIMILARITY: CONTAINS 1 CAND DOMAIN.
SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - Buropean Bloinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PEQLADAGEYYYGRNDDVKCFCCDGGLRCWESGDDPWIEHAKWFPRCE 327
                                                                                                                                                                                                                                                                                                                                                                            PF00653; BIR; 3.
PF00619; CARD; 1.
PF00097; zf-C3HC4; 1.
                                                                                                                                                                                                                                                                                                                SM00238; BIR; 3.
SM00114; CARD; 1.
SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genes."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                            : Zinc-finger;
27 94
167 233
253 320
444 512
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                                                                                         ξ
./b.,
1.
2; Repeat.
94 BIR 1.
233 BIR 2.
320 BIR 3.
CARD.
RING-TYPE.
AD7F73B6849317D1 CR
                                                                                      233
320
512
587
67198
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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.2e-26;
                                                                                      CRC64
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                   Length
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                        RESULT 6
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C. SUBCELULIAR LOCATION: CYTOPLASMIC (POTENTIAL).

C. TISSUE SPECIFICITY: EXPRESSED IN HEART, BRAIN, SPLEEN, LITURE, SKELETAL HUSCLE, KIDNEY, AND TESTIS.

C. SIMILARITY: BELONGS TO THE IAP FAMILY.

C. ISMILARITY: CONTAINS 3 BIR REPEATS.

C. SIMILARITY: CONTAINS 1 CAND DOMAIN.

C. ISMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 42; Conser
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                                   InterPro; IPR001370; BIR.
InterPro; IPR001315; CARD.
InterPro; IPR001841; Znf_ring.
InterPro; IPR001841; Znf_ring.
InterPro; IPR00185; BIR; 3.
Interpro; IPR00180; CARD; 1.
Interpro; IPR00180; INTERPRO; SMORT; SMO0114; CARD; 1.
SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
MEDILINE-95(128127; PubMed-95468107,
Rothe M., Pan M.-G., Henzel W.J., Ayres T.M., Goeddel D.V.;
"The THER2-TRAF signaling complex contains two novel proteins related
to baculoviral inhibitor of apoptosis proteins.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BIR3_MOUSE STANDARD; PRT; 612 AA. 062210; 009864; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 20-NUG-2001 (Rel. 40, Last annotation update) BACULOVIRAL LAP REPEAT CONTAINING PROTEIN 3 (
                                                                                                                                                                                                                                                                                                                  EMBL; U88909; AAC53532.1; MGD; MGI:1197009; Birc3.
                                                                                                                                                                                                                                                                                                                                                              EMBL; L49433; AAC42078.1; ..
                                                                                                                                                                                                                                                                                                                                                                                                         or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              between the Swiss Institute of Bioinf
the European Bioinformatics Institute
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROTEIN 2) (MIAP2) (MIAP-2).
BIRC3 OR API2 OR IAP2.
                   Apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      46:495-503(1997)
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                     -finger;
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272 QELASAGFYYTGHSDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 318
                                                                                            2e-25;
(INHIBITOR
                                                                                                                     1;
                                                                                                                     Indels
 õ
 APOPTOSIS
                                                                                                                     0
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Craniata; Vertebrata; |
Sciurognathi; Muridae;
                    Euteleostomi;
  Murinae; Mus
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SEQUENCE FROM N....
TISSUE-Skeletal muscle;
TISSUE-Skeletal muscle;
MEDLINE-98110590; Pubmed-9441758;
Liston P., Lefebvre C., Rong W.G., Xuan J.Y., Korneluk R.G.;
Liston P., Lefebvre C., Fong W.G., Xuan J.Y., Korneluk R.G.; protein

THE BIR MOTIFS REGION INTERACTS CTORS 1 AND 2 (TRAF1 AND TRAF2) ICH IS THEN RECRUITED TO THE TUM F2) TO TUMOR

SPLEEN, LUNG

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Best Local
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                                          InterPro. IPR001841: Znf_ring.
Pfam; PF00653; BIR; 2.
Pfam; PF00997; zf-C3HC4; 1.
SMART; SM00238; BIR; 2.
SMART; SM00238; BIR; 2.
SMART; SM00184; RING; 1.
PR0SITE; PS01282; BIR REPEAT_1; 2.
PR0SITE; PS0143; BIR_REPEAT_2; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1995 (Rel. 32, C
01-NOV-1995 (Rel. 32, L
20-AUG-2001 (Rel. 40, L
APOPTOSIS INHIBITOR 3 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REPEAT
REPEAT
REPEAT
DOMAIN
ZN_FING
CONFLICT
SEQUENCE
                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                  EMBL; L22564; AAB02610.1; -. EMBL; U75930; AAC59034.1; -.
                                                                                                                                                                            entitles requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                   Birnbaum M.J., Clem R.J., Miller L.K.;
"An apoptosis-inhibiting gene from a nuclear polyhedrosis virus encoding a polypeptide with Cys/His sequence motifs.";
J. Virol. 68:2521-2528(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Orgyia pseudotsugata multicapsid polyhedrosis virus (OpMNPV).
Viruses, dsDNA viruses, no RNA stage; Baculoviridae;
                       APOPTOS1S;
                                                                                                                               InterPro; IPR001370; BIR.
                                                                                                                                                                                                                                                                                                                                                 The
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                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                              sequence of the Orgyia pseudotsugata multinucleocapsid nuclear hedrosis virus genome.";
; Zinc-finger;
18 84
111 178
221 255
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177
262
447
565
380
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243 B
329 B
533 C
599 R
380 E
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87.5%;
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Last annotation update)
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BIR 1.
BIR 2.
RING-TYPE.
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BIR 2.
BIR 3.
CARD.
RING-TYPE.
RING-TYPE.
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Pred. No. 2.1e-25;
"" amatches 3;
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E08969D93C6C610D CRC64;
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RESULT
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Best Local
                                                                              REPEAT
ZN_FING
SEQUENCE
                                                                                                                                                              Pfam; PF00653; BIR; 2.
SMART; SM00238; BIR; 2.
SMART; SM00184; RING; 1
                                                                                                                                                                                                                                                                                                                                                 KANG W.K., CTOOK N.E., WINSTANLEY D., O'REILLY D.R.;
Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
-i-FUNCTION: ACTS BY BLOCKING CELLULAR APOPTOSIS RATHER
PREVENTING VIRAL STIMULATION OF APOPTOSIS.
-i- SIMILARITY: CONTAINS 2 BIR REPERTS.
-i- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "An apoptosis-inhibiting baculovirus motif.";
                                                                                                                                                                                                                                                                              use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
                                                                                                                                                                                                                                                                                                   This SMISS-PROT entry is copyright. It is produced through a c
between the Swiss Institute of Bioinformatics and the EMBL
the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=93188168; PubMed=8445726; Crook N.E., Clem R.J., Miller L.K
                                                                                                                 PROSITE; PS01282; BIR_REPEAT_1; 2.
PROSITE; PS50143; BIR_REPEAT_2; 2.
Apoptosis; Zinc-finger; Repeat.
REPEAT 7 73 BIR 1
                                                                                                                                                                                                InterPro; IPR001370; BIR. InterPro; IPR001841; Znf_ring.
                                                                                                                                                                                                                       EMBL; L05494; AAA43835.1; EMBL; U53466; AAB39098.1;
                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                   entities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-MEXICAN;
                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=28289;
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20-AUG-2001
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01-NOV-1995
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1 PEQLASAGEYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pomonella
                        29;
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30; Conserv
                                   Similarity
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275 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence u
annotation
                       ú
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                       Score 185; DB
Pred. No. 1.6e
5; Mismatches
                                                                             RING-TYPE. 84605448869CAD60 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 200; DB 1;
Pred. No. 2.3e-17;
6; Mismatches 12
                                                                                                       BIR 2.
                                                                                                                 BIR 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DF89175FDE85A708 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    update)
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                                                                                                                                                                                                                                                                 moved. Usage by and for (See http://www.isb-sib.
                                  DB 1;
.6e-15;
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                        13;
                                             Length
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126

PEQMADAGFFYTGYGDNTKCFYCDGGLKDWEPEDVPWEQHVRWFDRC

172

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RESULT 10
BIRG_MOUSE
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                                                                                                                                        Query Match
Best Local s
Matches 28
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OBJIE6: P81704: O09122: O09121:

O9JIE6: P81704: O09122: O09121:

OPAUG-2001 (Rel. 40, Created)

OPAUG-2001 (Rel. 40, Last sequence update)

OPAUG-2001 (Rel. 40, Last annotation update)

BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 1F (NE

INHIBITORY PROTEIN 6).

N BIRCIF OR NAIF6 OR NAIF-RS4.
 Q9JIB3;
                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss institute of Bioinformatics and the EWBL outst the European Bioinformatics institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed. Usage by and for commodified and this statement is not removed.
                BIRC_MOUSE
                                                                                                                                                                                                                                                                                                          MGD; MGI:1298222; Birclf.
InterPro; IPR001370; BIR.
Pfam; PF00653; BIR; 3.
SMART; SM00238; BIR; 3.
                                                                                                                                                                                                                                                                                                                                                                                                        entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Endrizzi M.G., Hadinoto V., Growney J.D. "Genomic sequence analysis of the mouse Genome Res. 10:1095-1102(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota: Metazoa; Chordata; Craniata; Vertebrata: Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI_TaxID-10090;
                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                              PROSITE; PS01282; BIR_REPEAT_1; PROSITE; PS50143; BIR_REPEAT_2;
                                                                                                                                                                                                                                                                                                                                                                EMBL; AF242431; AAF82751.1; -. EMBL; U66327; AAC52975.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genomics 38:405-417(1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              exon 5."
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Kunkel L.M., Dietrich W.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-97131520; PubMed-8975718;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-129/SVJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE OF 82-168 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-20414747; PubMed-10958627;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                    178
                                                                                                                                                                                                                                                                  Apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -1 - SIMILARITY: CONTAINS 3 BIR REPEATS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "The mouse region syntenic for human within the Lgnl critical interval and
                                                                                                            1 PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: PREVENTS MOTOR-NEURON
                                                                                  PRALSAAGFVFTGKRDTVQCFSCGGSLGNWEEGDDPWKEHAKWFPKCE
                                                                                                                                        Similarity 58.
                                                                                                                                                                                                             1403
                                                                                                                                                                                                                                                                  Repeat; Multigene
                STANDARD;
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                                                                                                                                                    61.68;
58.38;
                                                                                                                                                                                                             159823 MW;
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                                                                                                                                                                                                                                     _1; 2.
_2; 3.
family.
BIR 1.
BIR 2.
                                                                                                                                       Score 181; DB 1;
Pred. No. 2.2e-14;
5; Mismatches 14
                                                                                                                                                                                                                            BIR
              PRT;
                                                                                                                                                                                                             9D4912503358C4E9 CRC64;
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                1402 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                s.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene array.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NEURONAL APOPTOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INDUCED BY
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                                                                                                                                                                   Length 1403;
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inle copies of !
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BIRA_MOUSE
IDA_MOUSE
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AC QQWKS
DT 20-AUG
DT 20-AUG
DT 20-AUG
DE BACULO
DE INHIBI
GN BIRCIA
OS MUS MUS MUS CLU
OC EUKARY
CC MEMMAT
OX NCBLT
RN [1]
RP SEQUEN
RA YEARSH
RT COLONI
RT COLONI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT
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Best Local
                                                                                                                                                                                                                                                                             0900K5; 098017; 09J1B5;
20-AUG-2001 (Rel. 40, Created)
20-AUG-2001 (Rel. 40, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation updat)
BACULOVIRAL IAP REPEAT—CONTAINING PROTEIN 1
INHIBITORY PROTEIN 1).
                                          "Clonday and characterization of the multiple copies of th homologue of NAIP (neuronal apoptosis inhibitory protein). Submitted (JUN-1997) to the EMBL/GenBank/DDBJ databases.
  SEQUENCE FROM N.A
                                                                                                                           SEQUENCE FROM N.A
                                                                                                                                                                      NCBI_TaxID=10090;
                                                                                                                                                                                          Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                     Mus musculus (Mouse).
                                                                                                                                                                                                                                                           BIRCIA OR NAIPI OR NAIP.
                                                                                                                                                                                                                                                                                                                                                                                                                BIRA_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS01282; BIR_REPEAT_1; 2. PROSITE; PS50143; BIR_REPEAT_2; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MGD; MGI:1858256; Birclg.
InterPro; IPR001370; BIR.
Pfam; PF00653; BIR; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; AF242433; AAF82749.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome Res.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Endrizzi M.G., Hadinoto V.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse)
Eukaryota; Metazoa; (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BIRC1G OR NAIP7
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20-AUG-2001 (Rel. 40, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 1G
INHIBITORY PROTEIN 7).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=20414747; PubMed=10958627;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID-10090;
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                                                                                                         araghi 2.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   poptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     178 PRVLSAAGFVFTGKRDTVQCFSCGGSLGNWEEGDDPWKEHAKWFPKCE 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: CONTAINS 3 BIR REPEATS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION: PREVENTS MOTOR-NEURON APOPTOSIS
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28; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           M.G., Hadinoto V., Growney J.D., Miller we sequence analysis of the mouse Naip genees, 10:1095-1102(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (; Repeat; Multigene (
60 127
159 227
278 345
1402 AA; 159662 My
                                                                                                         Korneluk R.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          58.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 BIR 3.
159662 MW; C1DFFBA359893E0D CRC64;
                                                                                                         MacKenzie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     family.
BIR 1.
BIR 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 180; DB 1;
Pred. No. 2.9e-14;
6; Mismatches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                          Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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PROTEIN 1A (NEURONAL
                                                                                                                                                                                                                                                                                                                                                                                                                1403
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                                                                                                       A.E.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14;
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BIRE_MOUSE
ID BIRE_MO
BIRE_MO
AC Q9R016
AC Q9R016
DT 20-AUG
DT 20-AUG
DT BACULO
DE BACULO
GN BIRCIE
GN BIRCIE
GN BIRCIE
GN MUS mus
OC Musman
OC MAMMAN
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Best Loc
Matches
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EMBL; AF13491; AAD56763.1; -.
EMBL; AF24432; AAF82752.1; -.
MGD; MGI:1298223; Bircla.
InterPro; IPR001370; BIR.
Pfam; PF00653; BIR; 3.
SMART; SM00238; BIR; 3.
                                                                                                                                                                                                          BIRE MOUSE STANDARD; PRT; 1403 AA. 09R015; 098029; P81703; 009122; 009121; 20-AUG-2001 (Rel. 40, Created) 20-AUG-2001 (Rel. 40, Last sequence update) 20-AUG-2001 (Rel. 40, Last annotation update) BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 1E
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REPEAT
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SEQUENCE FROM N.A
                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                  INHIBITORY PROTEIN 5).
BIRC1E OR NAIP5 OR NAIP-RS3
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apoptosis;
REPEAT
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MEDLINE-20414747;
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                                                   NCBI_TaxID-10090;
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                                                                                  Eutheria;
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58.3%;
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                                                                               Sciurognathi;
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                                                                                  Muridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 1403;
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                                                                                  Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-99417674; PubMed-10486205; Endrizzi M., Huang S., Scharf J.M., Ke Kunkel L.M., Miller W., Dietrich W.F.; "Comparative sequence analysis of the interval.";
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Mamm. Genome 10:1032-1035(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF131205; AAD56760.1;
EMBL; U66326; AAC52974.1; -.
MGD; MGI:1298220; Bircle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; AF135492; AAD56764.1; -. EMBL; AF131205; AAD56760.1; -.
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Kunkel L.M., Dietrich W.F.;
"The mouse region syntenic for human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=129/SVJ;
MEDLINE=97131520; PubMed=8975718;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00653; BIR; 3
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1. 2. 3. 6 mily. Family. BIR 1. BIR 2. BIR 2. S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S > R S 
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Matches
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Best Local
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PRVLSAAGFVFTGKRDTVQCFSCGGSLGNWEEGDDPWKEHAKWFPKCE 225
                       PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 48
                                                               Similarity
                                                 61.2%;
ilarity 58.3%;
Conservative
                                                 Score 180; DB
Pred. No. 2.9e
6; Mismatches
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CONFLICT CONFLICT

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521 533 538 538 1092 1129 1127 1137

REPEAT

SEQUENCE

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159695

B27F645043BCEC42 CRC64;

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BIRA_MOUSE
ID BIRA_MOUSE
AC Q6099; OOB
AC C6099; OOB
AC C6
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Pfam; PF00053; BIR; 3.
SMART; SM00238; BIR; 3.
SMART; SM00184; RING; 1.
SM0184; RING; 1.
SM0184; BIR_REPAT_1; 3.
PROSITE; PS0143; BIR_REPEAT_2; 3.
Apoptca1s; Zinc-finger; Repeat.
REPEAT 26 93 BIR 1.
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01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
01-AUG-2001 (Rel. 40, Last annocation update)
20-AUG-2001 (Rel. 40, Last annocation update)
BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 4 (INHIBITOR OF APOPTOSIS PROTEIN 3) (X-LINKED INHIBITOR OF APOPTOSIS PROTEIN) (X-LINKED IAP)
(IAP HOMOLOG A) (MIAP3) (MIAF-3).
BIRC4 OR API3 OR XIAP OR AIFA OR MIHA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-C57BL/6 x CBA; TISSUE-Liver;
MEDLINE-96209843; PubMed-8643514;
Uren A.G., Pakusch M., Hawkins C.J., Puls K.L., Vaux D.L.;
"Cloning and expression of apoptosis inhibitory protein homologs that
function to inhibit apoptosis and/or bind tumor necrosis factor
receptor-associated factors.";
record receptor-associated factors.";
Proc. Natl. Acad. Sci. U.S.A. 93:4974-4978(1996).
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR001370; BIR.
InterPro; IPR001841; Znf_ring.
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BIR4_RAT
ID BIR4_RAT
AC Q9R0I6;
DT 20-AUG-2001
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ZN_FING
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-AUG-2001 (Rel. 40, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 4 (INHIBITOR OF APOPTOSIS PROTEIN 3) (X-LINKED INHIBITOR OF APOPTOSIS PROTEIN) (X-LINKED IAP)
(IAP HONOLOG (RIAP3) (RIAP3).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Rattus norvegicus X-linked inhibitor of apoptosis (riap3) mRNA. Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: APOPTOTIC SUPPRESSOR. INHIBITOR OF CASPASE-3 AND CASPASE-7 (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: CYTOPLASMIC (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                             the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on the EMBL outstation the Buropean Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus norvegicus (Rat),
Eukaryota: Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                                                                                   REPEAT
                                                                                                                                                                                                APOPTOSIS;
REPEAT
                                                                                                                                                                                                                               PROSITE;
                                                                                                                                                                                                                                             PROSITE;
                                                                                                                                                                                                                                                            SMART; SM00238; BIR; 3.
SMART; SM00184; RING; 1
                                                                                                                                                                                                                                                                                       Pfam; PF00053; BIR; 3.
Pfam; PF00097; zf-C3HC4; 1.
                                                                                                                                                                                                                                                                                                                    InterPro; IPR001370; BIR. InterPro; IPR001841; Znf_
                                                                                                                                                                                                                                                                                                                                                 EMBL; AB033366; BAA85304.1; -.
                                                                                                                                                                                                                                                                                                                                                                                or send an email to license@isb-slb.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -i- SIMILARITY: BELONGS TO THE IAP FAMILY.
-i- SIMILARITY: CONTAINS 3 BIR REPEATS.
-i- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Saito
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10116;
 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  281 EQLARAGEYALGEGDKVKCFHCGGGLTDWKPSEDPWEQHAKWYPGCK
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nes 29; Conser
                  EQLARAGEYALGEGDKVKCFHCGGGLTDWKPSEDPWEQHAKWYPGCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29;
                                                                                                                                                                                                                             PS50143; BIR_REPEAT_2;
                                                                                                                                                                                                                                            PS01282; BIR_REPEAT_1;
                                                                                                                                  264
449
496
                                                                                                                                                                                                Zinc-finger; Repeat.
26 93 BJ
                                                           Conservative
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56072 MW;
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BIR 2.
BIR 3.
RING-TYPE.
RING-TYPE.
RYM; E250E3C77461A469 CRC64;
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Pred. No. 2.6e-14;
                                                           Pred. No. 2.66
5; Mismatches
                                                                          Score 177; DB 1;
Pred. No. 2.6e-14;
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BIR4_HUMAN ID BIR4_F

BIR4_HUMAN

STANDARD

PRT;

497

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RESULT 15

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EMBL, ALIZ1601; CAB95312.1; --
MIN; 300079; --
InterPro; IRR001370; BIR,
InterPro; IRR001841; Znf_ring,
Pfam; PF00053; BIR; 3
Pfam; PF00097; zf-C3HG4; 1.
SMART; SM00184; RING; 1.
SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-9626286; PubMed-8654366;
Duckett C.S., Nava V.E., Gedrich R.W., Clem R.J., van Dongen J.L., Gilfillan M.C., Shiels H., Hardwick J.M., Thompson C.B.;
"A conserved family of cellular genes related to the baculovirus iap gene and encoding apoptosis inhibitors.";
EMBO J. 15:2685-2694(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
TISSUE-Fetal brain;
MEDLINE-96149249; PubMed-8552191;
MEDLINE-96149249; PubMed-8552191;
Liston P., Roy N., Tamai K., Lefebvre C., Baird S., Cherton-Horvat Liston P., Roy N., Tamai K., Lefebvre C., Baird S., Cherton-Horvat Farabani R., McLean M., Ikeda J., Mackenzle A., Korneluk R.G.;
"Suppression of apoptosis in mammalian cells by NAIP and a related family of LAP genes."
[antly of LAP genes."]
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01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
01-0CT-1996 (Rel. 34, Last sequence update)
02-AUG-2001 (Rel. 40, Last annotation update)
BACULOVIRAL TAR REPAIR-COMPAINING PROFEIN 4 (INHIBITOR OF APOPTOSIS
BACULOVIRAL TAR REPAIR-COMPAINING PROPEIN 4 (INHIBITOR OF APOPTOSIS PROTEIN) (X-LINKED INHIBITOR OF APOPTOSIS PROTEIN) (X-LINKED IAP)
BROYEL N (HILP).
LIRCH PROFEIN (HILP).
BIRC4 OR API3 OR TAP3 OR XIAP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its grown profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Elikaryota: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi;
Mammalia: Eutheria: Primates: Catarrhini; Hominidae: Homo.
MCBL_TaxID-9606;
                                                                                                                                PROSITE; PS01282; BIR_REPEAT_1; PROSITE; PS50143; BIR_REPEAT_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; U45880;
EMBL; U32974;
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- I SIMILARITY: BELONGS TO THE IAP EAMILY.

- I SIMILARITY: CONTAINS 3 BIR REPEATS.

- I SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Deveraux O.L., Takahashi R., Salvesen G.S., Reed J.C.; "X-linked IAP is a direct inhibitor of cell-death prot Nature 388:300-304(1997).

-i- FUNCTION: APOPTOTIC SUPPRESSOR. INHIBITOR OF CASPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-97373959; PubMed-9230442;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: APOPTOTIC SUPPRESSOR. INHIBITOR OF CASPASE. 7.
SUBCELLULAR LOCATION: CYTOPLASMIC.
TISSUE SPECIFICITY: UBIQUITOUS, EXCEPT PERIPRESSOR.
26 93
163 230
265 330
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AAC50518.1; ~.
                                                                                                   Repeat;
    BIR
BIR
                                                                                                   Thiol protease inhibitor.
    ω Ν ⊢
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CONFLICT
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Q -> P (IN
                                                                        5 -> C (IN REF. 1).

-> P (IN REF. 2).

9D394C16D45EB635
                                        DB 1;
                                 13;
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Search completed: January 7, 2002, 16:05:26 Job time: 1405 sec